

GenCore version 6.3  
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:11:25 ; Search time 22 Seconds  
(without alignments)  
28686.456 Million cell

updates/sec

Title: US-09-300-482-27  
Perfect score: 229  
Sequence: 1  
cagaccttattttttctgtc.....cttcaatttatggggaacaa 229

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 2888648 seqs, 1391227531 residues

Total number of hits satisfying chosen parameters: 5777296

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Pending\_Patents\_NA\_New:\*  
1:  
/ABSS/Data/CRF/ptodata/2/pna/PCT\_NEW\_COMB.seq:\*  
2:  
/ABSS/Data/CRF/ptodata/2/pna/US08\_NEW\_COMB.seq:\*  
3:  
/ABSS/Data/CRF/ptodata/2/pna/US09\_NEW\_COMB.seq:\*  
4:  
/ABSS/Data/CRF/ptodata/2/pna/US10\_NEW\_COMB.seq:\*  
5:  
/ABSS/Data/CRF/ptodata/2/pna/US11\_NEW\_COMB.seq:\*  
6:  
/ABSS/Data/CRF/ptodata/2/pna/US12\_NEW\_COMB.seq:\*  
7:  
/ABSS/Data/CRF/ptodata/2/pna/US12\_NEW\_COMB.seq1:\*  
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